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=> d his

     (FILE 'HOME' ENTERED AT 15:50:06 ON 16 JAN 2009)

     FILE 'MEDLINE, EMBASE, BIOSIS, SCISEARCH, DISSABS, TOXCENTER' ENTERED AT
     15:51:09 ON 16 JAN 2009
L1       2729 S (HEINRICH M?/AU) OR (CORLESS C?/AU)
L2      10611 S (FLETCHER J?/AU) OR (DEMETRI G?/AU)
L3      13149 S L1 OR L2
L4        218 S L3 AND (PDGF?)
L5          0 S L3 AND CD140?
L6        180 S L3 AND (PLATELET(1A)DERIVED)
L7        264 S L4 OR L6
L8        207 S L7 AND MUTAT?
L9        178 S L8 AND (TUMOR? OR CANCER? OR CARCIN?)
L10       109 S L8 AND NEOPLAS?
L11       186 S L9 OR L10
L12       456 S L3 AND GIST
L13       128 S L4 AND GIST
L14        78 DUP REM L11 (108 DUPLICATES REMOVED)
L15         7 S L14 AND PY<=2002
L16       199 DUP REM L12 (257 DUPLICATES REMOVED)
L17        26 S L16 AND PY<=2002
L18        31 S L15 OR L17
L19        31 DUP REM L18 (0 DUPLICATES REMOVED)
L20      2371 S (PDGFRA)
L21        99 S (PDGFR-A)
L22      2344 S (PDGFR(2A)ALPHA)
L23       327 S (PDGF-R(2A)ALPHA)
L24      2748 S (PDGF(2A)RECEPTOR(2A)ALPHA)
L25      4759 S (PLATELET(2A)DERIVED(5A)RECEPTOR(10A)ALPHA)
L26        23 S CD140A
L27      8628 S L20 OR L21 OR L22 OR L23 OR L24 OR L25 OR L26
L28      2386 S L27 AND MUTAT?
L29         0 S L27 AND (2072 OR 2073 OR 2074 OR 2075)
L30         1 S L27 AND (2076 OR 2077 OR 2078 OR 2079 OR 2080)
L31         0 S L27 AND (2081 OR2082 OR 2083 OR 2084 OR 2085)
L32         0 S L27 AND (2086 OR 2087 OR 2088 OR 2089 OR 2090)
L33         0 S L27 AND (2091 OR 2092 OR 2093 OR 2094 OR 2095)
L34         0 S L27 AND (2096 OR 2097 OR 2098 OR 2099 OR 2100)
L35         0 S L27 AND (2101 OR 2102 OR 2103 OR 2104 OR 2105 OR 2106 OR 210
L36         0 S L27 AND (2916 OR 2917 OR 2918 OR 2919 OR 2920 OR 2921 OR 292
L37         0 S L27 AND (2923 OR 2924 OR 2925 OR 2926 OR 2927 OR 2928 OR 292
L38         0 S L27 AND (2930 OR 2931 OR 2932 OR 2933 OR 2934 OR 2935 OR 293
L39        39 S L27 AND (560 OR 561 OR 562 OR 563 OR 564 OR 565)
L40       156 S L27 AND (566 OR 567 OR 568 OR 569 OR 570 OR 571)
L41       186 S L39 OR L40
L42       117 DUP REM L41 (69 DUPLICATES REMOVED)
L43        48 S L27 AND (841 OR 842 OR 843 OR 844 OR 845 OR 846 OR 847 OR 84
L44       161 S L42 OR L43
L45       127 DUP REM L44 (34 DUPLICATES REMOVED)
L46        70 S L45 AND MUTAT?
L47        27 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(560 OR 561 OR 562
L48        11 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(566 OR 567 OR 568
L49        37 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(841 OR 842 OR 843
L50        16 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(845 OR 846 OR 847
L51        60 S L47 OR L48 OR L49 OR L50
L52        20 DUP REM L51 (40 DUPLICATES REMOVED)
L53         0 S L52 AND PY<=2002
L54         2 S L52 AND PY<=2003
L55       349 S L27 AND ((EXON(1A)10)OR(EXON(1A)12))
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L56         111 DUP REM L55 (238 DUPLICATES REMOVED)
L57           1 S L56 AND PY<=2002
L58         186 S L27 AND ((EXON(1A)18))
L59          67 DUP REM L58 (119 DUPLICATES REMOVED)
L60           0 S L59 AND PY<=2002

=> log y
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